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FIGURE 1

420	425	430	435	440	445	450	455	460	465	470	475	480	485	490	495
A	AAG	GGG	CGA	ATT	CAG	G	ATG	CCG	TGC	AGC	AGA	GGT	CTG	ATG	CTC
T	TTC	CCC	CCG	TAA	GTC	C	TAC	GGC	ACG	TGG	TCT	CGA	GAC	TAA	GAG
Met Pro Cys Ser Arg Ala Leu Ile Leu Gly Val Leu Ala Leu Asn Thr Met Leu Ser Leu>															
EcoRI															
500	505	510	515	520	525	530	535	540	545	550	555	560	565	570	
TGC	CGA	GGT	GNA	GAC	GAC	GAC	GAC	GAC	GAC	GAC	GAC	GAC	GAC	GAC	ATT
ACG	CCG	CCA	CTT	CAG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	TAA
Cys	Gly	Gly	Glu	Asp	Asp	Ile	Glu	Ala	Ala	Asp	Ile	Val	Gly	Pro	Gly
Ile>															
575	580	585	590	595	600	605	610	615	620	625	630	635	640	645	650
GGC	CAG	TAC	ACA	CAT	GAA	TAT	GAT	GAT	GAT	GAT	GAT	GAT	GAT	GAT	CTT
CCG	GTC	ATG	TGT	GTA	CTT	AAA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	GNA
Gly	Glu	Thr	His	Glu	Pro	Asp	Gly	Asp	Glu	Leu	Pro	Thr	Val	Arg	Leu>
655	660	665	670	675	680	685	690	695	700	705	710	715	720	725	730
CCG	GAG	TAT	ACC	CTA	ACT	ACC	ACT	ACT	ACT	ACT	ACT	ACT	ACT	ACT	CTT
CGA	CTC	AAA	CCG	CTT	ACC	TAT	GAG	AAA	CTC	CTG	CTG	CTG	CTG	CTG	CTT
Pro	Glu	Pro	Glu	Glu	Pro	Glu	Gly	Gly	Leu	Asn	Ile	Ala	Glu	Lys	His
735	740	745	750	755	760	765	770	775	780	785	790	795	800	805	
ATC	TGC	ACT	AAG	ACG	TCA	ACT	ACC	ACT	ACC	ACT	ACC	ACT	ACC	ACT	CTT
TAG	AAC	TCA	TTC	ACT	TTA	ACT	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	GAG
Ile	Leu	Thr	Lys	Arg	Ser	Pro	Ala	Thr	Asn	Glu	Ala	Pro	Val	Pro	Val>
810	815	820	825	830	835	840	845	850	855	860	865	870	875	880	885
CTG	CTG	CTT	CAG	CCC	AAC	ACC	CTT	ATC	TCC	TTC	GTC	GTC	GTC	GTC	AGA
GAC	GAC	CCA	CTC	GGG	TTC	TGG	GNA	TAG	ACC	AAA	CAC	CTG	TAG	CTG	CTT
Leu	Leu	Gly	Glu	Pro	Asn	Thr	Leu	Ile	Cys	Pro	Val	Asp	Asn	Ile	Thr
890	895	900	905	910	915	920	925	930	935	940	945	950	955	960	
ATC	ACC	AAG	TCA	GTC	ACA	GAC	GGC	GTT	TAT	GAG	ACC	AGC	TTC	CTC	CTT
TTC	TGC	TTC	ACT	CAG	TGT	CTG	CCG	CNA	ATA	CTC	TGC	NAG	CAG	TTC	AGA
Asn	Ser	Lys	Ser	Val	Thr	Asp	Gly	Val	Thr	Glu	Thr	Ser	His	Lys	Ser>

[illegible]

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FIGURE 2

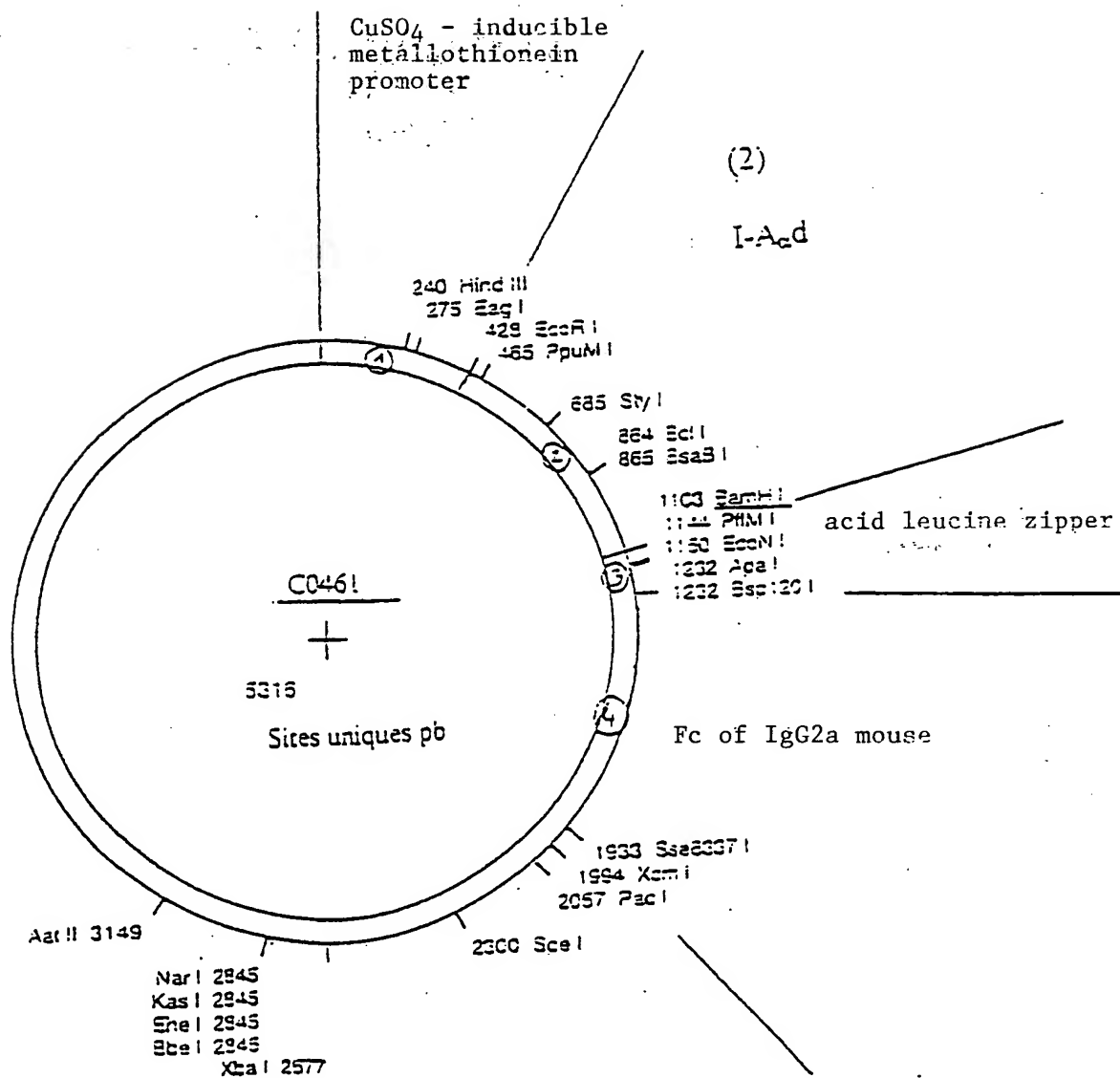


FIGURE 3

[illegible]

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965 970 975 980 985 990 995 1000 1005 1010 1015 1020 1025 1030 1035 1040
 ACT CIG GIC TGT TCG GTG ACA GAT TFC TAC CCA CCC AAG ATC AAA GTC CGC TCG TTC AGG ANT GCC CAG GAG ACA
 Thr Leu Val Cys Ser Val Thr Asp Phe Tyr Pro Ala Lys Ile Lys Val Arg Trp Phe Arg Asn Gly Gln Glu Thr>

 1045 1050 1055 1060 1065 1070 1075 1080 1085 1090 1095 1100 1105 1110 1115 1120
 GTC CGC GTC TCA TCC ACA CAG CTT ATT ACC ANT CCG GAC TCG ACC TTC CAG GTC CIG GTC ATG CIG GAG ATG ACC CCT
 Val Gly Val Ser Ser Thr Gln Leu Ile Arg Asn Gly Asp Trp Thr Phe Gln Val Leu Val Met Leu Glu Met Thr Pro>

 1125 1130 1135 1140 1145 1150 1155 1160 1165 1170 1175 1180 1185 1190 1195
 CMT CAG GCA GAG GTC TAC ACC TCC CMT CIG CAG CMT CCC ACC CIG AAG AGC CCC ATC ACT CIG GAG TCG AGG GCA CAG
 His Gln Gly Glu Val Tyr Thr Cys His Val Glu His Pro Ser Leu Lys Ser Pro Ile Thr Val Glu Trp Arg Ala Gln>

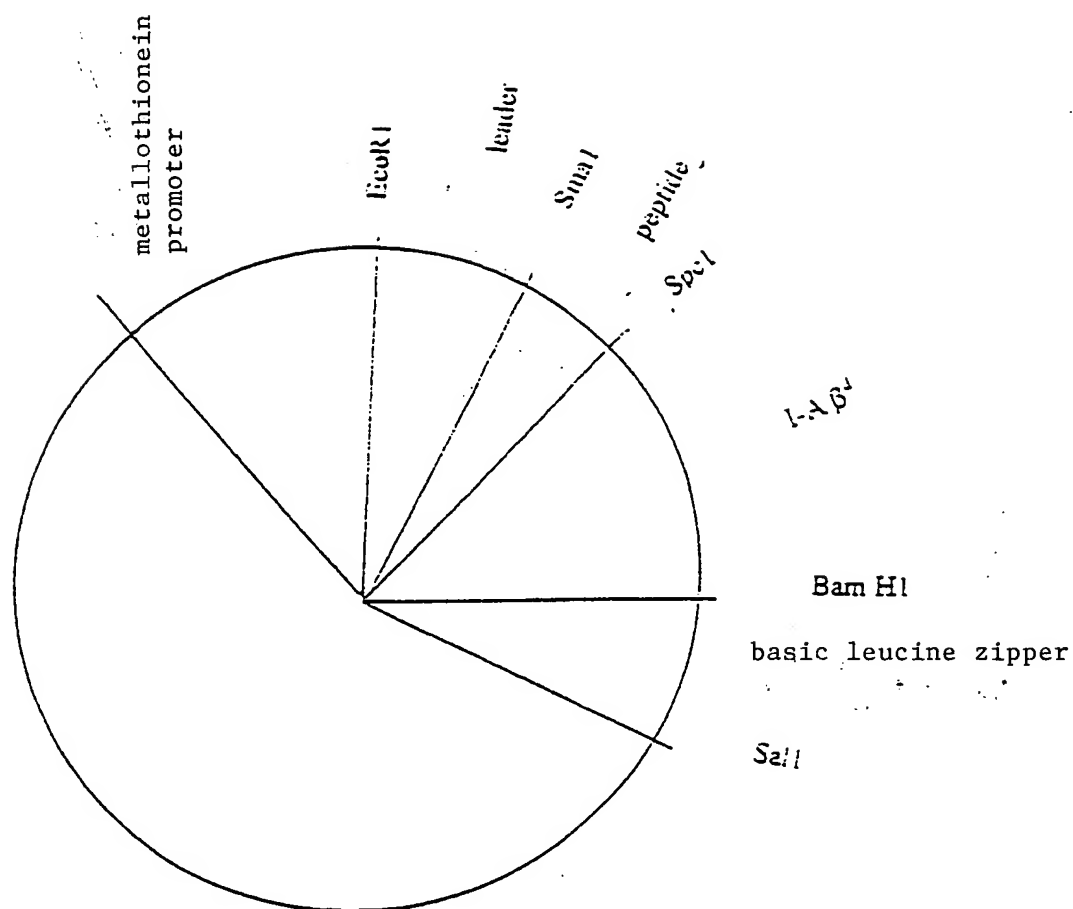
 1200 1205 1210 1215 1220 1225 1230 1235 1240 1245 1250 1255 1260 1265 1270 1275
 TCC GAG TCT GCC CCG ACC AAG CCA GGT CCA TCA TCC ACT ACA GGT CCA TCA CCF CAG TIG AAA AAG AAA TIG CAA GCA
 Ser Glu Ser Ala Arg Ser Lys Gly Gly Gly Gly Ser Thr Thr Ala Pro Ser Ala Gln Leu Lys Lys Lys Leu Glu Ala>
 Linker
 1280 1285 1290 1295 1300 1305 1310 1315 1320 1325 1330 1335 1340 1345 1350
 CIG AAG AAA AAG AAC GCT CAG CIG AAG TCG AAA CTT CAA GCC CIG AAG AAG AAA CIG CCC CAG CMT CMT CMT CMT CMT
 Leu Lys Lys Lys Asn Ala Gln Leu Lys Trp Lys Leu Glu Ala Leu Lys Lys Lys Leu Ala Gln His His His His
 basic leucine zipper
 1355 1360 1365 1370
 CMT TCA GT CCA CCF CC
 His ***> Salt
 HISTAG

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FIGURE 4



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FIGURE 5

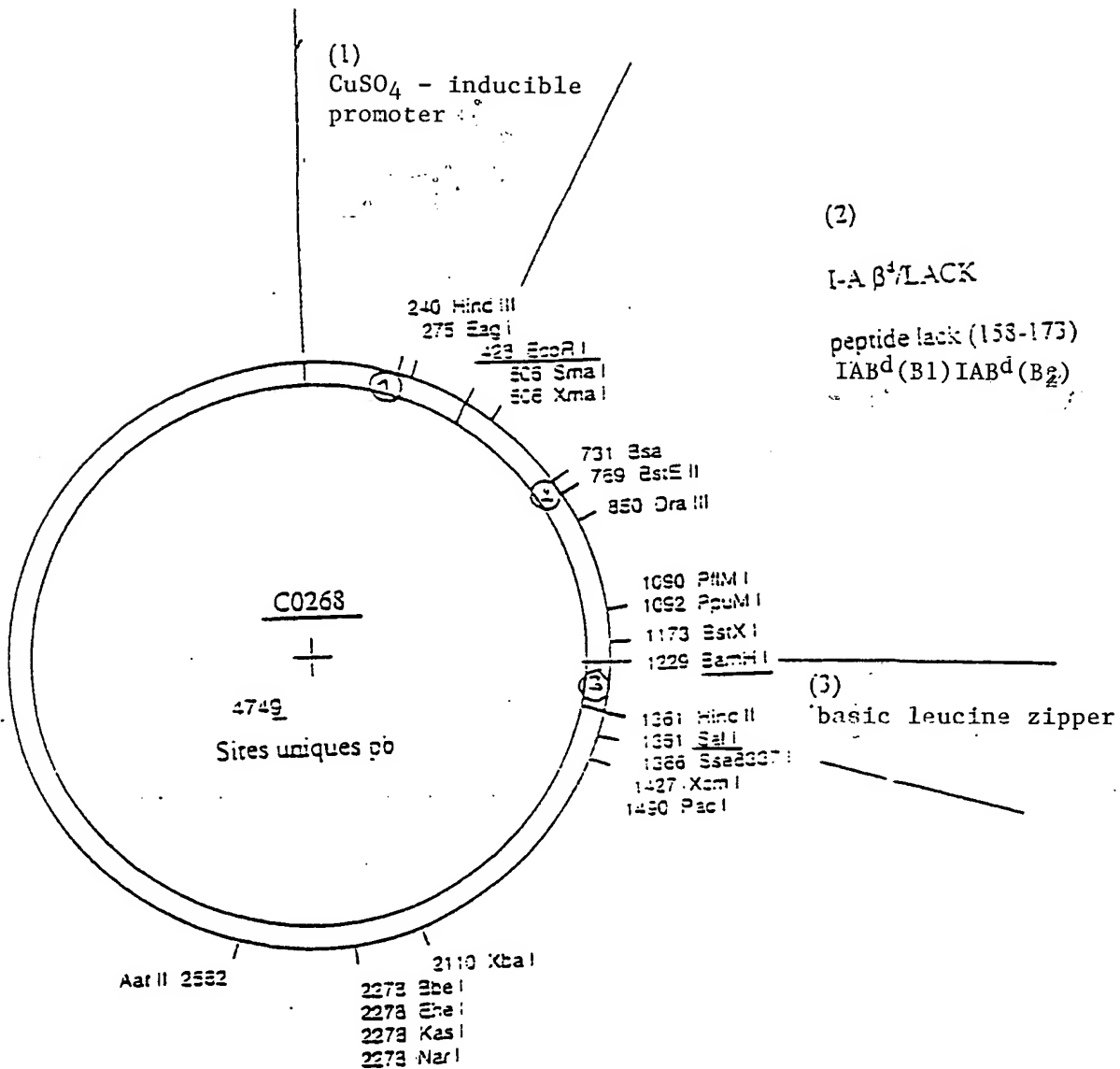


FIGURE 6

